

**AMENDMENTS TO THE CLAIMS**

This listing of claims will replace all prior versions and listings of claims in the application:

**Listing of claims:**

1. (currently amended) A method for SNP typing which comprises the steps of:  
  
simultaneously amplifying a plurality of nucleotide sequences comprising at least one or more sites of single nucleotide polymorphism using genomic DNA whose amount is 10-40 ng per 100 sites and a plurality of primer pairs; and  
  
typing for distinguishing the site(s) of single nucleotide polymorphism of ~~nucleotides contained in a plurality of~~ nucleotide sequences amplified in the above amplification step using the amplified nucleotide sequences.
  
2. (currently amended) The method for SNP typing according to claim 1, wherein said step of amplifying employs a ~~the~~ polymerase chain reaction using a hot start method.
  
3. (original) The method for SNP typing according to claim 1, wherein said step of amplifying employs 50 pairs or more primers.

4. (currently amended) The method for SNP typing according to claim 1, wherein said step of typing employs an ~~Invader~~ INVADER assay or a ~~TagMan~~ TAQMAN PCR method.

5. (currently amended) The method for SNP typing according to claim 1, wherein amplification is carried out on from 10 ng to 40 ng of genomic DNA.